us-10-676-2

Applicant Con

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Q10283 schizosacch
P2524 bacteriopha
Q8tym methanopyru
933400 saccharomyr
208493 home sapien
P14376 escherichia
P97793 mus musculu
Q10187 schariosacch
P3323 escherichia
Q7vn67 haemophilus
attus norv
       helicobacte
SSA1_RAT
GATA_HELPY
HMDH_SCHPO
VGA_BPAL3
                               SYFA METKA
R101 FEAST
CA4C HUMAN
RCSC ECOLI
ALK MOUSE
VAWD SCHPO
LLDR ECOLI
PPG HAEDU
61.55
61.55
61.55
61.55
61.55
61.55
61.55
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ALIGNMENTS

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MEDLINE=9107054; PubMed=8416957; MEDLINE=91107054; PubMed=8416957; MEDLINE=91107054; PubMed=8416957; Mild J. Rosen B.P.; PubMed=1107054112 PubMed=87258(1993).

-1. FUNCTION: TRANSCRIPTIONAL REPRESSOR FOR THE ARS OPERON. ARSR IS EXPRESSION. THE REPRESSIVE BFFECT OF ARSR IS ALLEVIATED BY OXYIONS OF +111 OXIDATION STATE OF ARSENIC, ANTIMONY, AND BISMUTH, AS WELLASUBMITE. Binds DNA as a homodimer.
-1. SUBUNIT: Binds DNA as a homodimer.
-1. SIBILARITY: BELONGS TO THE ARSR FAMILY OF TRANSCRIPTIONAL
                                                                                                                                                                                                              MEDLINE-90174986; PubMed-2408017; San Prancisco M.J.D., Hope C.L., Cwolabi J.B., Tisa L.S., Rosen B.P.; "Identification of the metallorgulatory element of the plasmid-encoded arsenical resistance operon."; Nucleic Acids Res. 18:619-624(1990).
                                                                                                                                           Bacteria, Proteobacteria, Gammaproteobacteria, Enterobacteriales,
Enterobacteriaceae, Escherichia.
NCBL_TaxID=562;
                                                                                                                                                                                                                                                                                                                   MEDLINE=92157659; PubMed=1838573; Wu J., Rosen B.P.; Wu T. Kosen B.P.; With Arbar protein is a trans-acting regulatory protein."; Mol. Microbiol. 5:1331-1336(1991).
                                                     01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
                             PRT; 117 AA
                                                                                                Arsenical resistance operon repressor
                             STANDARD;
                                                                                                                                                                                                            SEQUENCE PROM N.A.
                                                                                                                        Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGULATORS
                                                                                                                                         Plasmid R773
                           ARR1 ECOLI
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HSSP; P30340; 1SMT.
InterPro; IPR001845; HTH ArsR.
Pfam; PR01022; HTH 5; 1.
PRINTS; PR00778; HTHARSR.

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PROSITE; PS00846; HTH ARSR FAMILY; 1.
Plasmid; Arsenical resistance; Transcription regulation; Repressor;
PNA-binding.
DNA-binding.
33 52 H-T-H MOTIF (POTENTIAL).
SEQÜENCE 117.AA; 13198 MW; 1F0010766E4FD886 CRC64; fuical resistance, Transcription regulation, Repressor, Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Escherichia. NCBI\_TaxID=562; . 0 AS ARSENATE (AS(V)).
SUBUNIT: Binds DNA as a homodimer.
SIMILARITY: BELONGS TO THE ARS FAMILY OF TRANSCRIPTIONAL 100.0%; Score 506, DB 1; Length 117; 100.0%; Pred. No. 2e-48; tive 0; Mismatches 0; Indels 33 52 H-T-H MOTIF (POTENTIAL). 117 AA; 13198 MW; 1F0D10766E4FD886 CRC64; 61 LDRKQGKWVHYRLSPHIPSWAAQIIEQAWLSQQDDVQ 97 LDRKQGKWVHYRLSPHIPSWAAQIIEQAWLSQQDDVQ 97 01-OCT-1996 (Rel. 34, Created) 01-OCT-1996 (Rel. 34, Last sequence update) 15-DEC-1998 (Rel. 37, Last annotation update) Arsenical resistance operon repressor. FAMILY; 1 HTH\_ArsR HTH ARSR, 1. Query Match
Best Local Similarity 100.7
Matches 97; Conservative STANDARD; EMEL; U38947; AAB09624.
HSSP; P30340; 1SMT.
INCEPPRO; IPR001849; HTP
Pfam; PF01022; HTP 5; 1
PR.HTMS; PR00778; HTHARS;
SMART; SM00418 HTHARS; Escherichia coli. Plasmid IncN R46. PROSTE; PSO094 Plasmid, Arshni DNA-binding DNA BIND 33 SEQUENCE 117 REGULATORS ARR2\_ECOLI P52144; ARR2\_BCOLI RESULT 2 0.000 H H H H R 8 0 0 0 F STAAU

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10-0CT 品品資訊品 õ g ð g 당 Š CADC. Staphy Bacter NCBI 7 [1] RESULT YD25\_ ID AC DT DT 8 Query Best Match cadmi TRANS indus genes This betw

88.3%; Score 447; DB 1; Length 117; ilarity 87.6%; Pred. No. 5.7e-42; Conservative 5; Mismatches 7; Indels 85; est Local hu Jun

Similarity

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atches

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Gaps

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7 33 52 H-T-H MOTIF (POTENTIAL). 117 AA; 12999 MW; 4E2D132F1F011AF6 CRC64;